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Compound

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Advanced Keyword Query for: dipeptidyl peptidase IV

♠ 1234567

Protyl Tripeptidyl Aminopeptidase Complexed with an Inhibitor

Release Date: 19-Feb-2008 Exp. Method: Characteristics Resolution: 2.20 Å

Hydrolase Classification Polymer: 1 Molecule: Dipeptidyl aminop

no.: 3.4.14.-Xu. Y., Nakajima, Y., Ito. K., Yoshi Authors

Human Dipeptidyl Peptidase IV in complex with a Triazolopiperazine-

putative Fragment residues 39-732 (

based beta amino acid inhibitor Release Date: 06-Nov-2007 Exp. Method.

Characteristics Resolution: 2.30 Å

Hydrolase Classification

Polymer: 1 Molecule: Dipeptidyl peptida Fragment: Catalytic Domain Mutation: S Compound

Chains: A,B EC no.: 3.4.14.5 59 Polymer: 2 Molecule: SUGAR (2-MER)

Polymer: 3 Molecule: SUGAR (2-MER)

Scapin, G. Authors

Classification

Human dipeptidyl peptidase iv/ cd26 in complex with a 4-aryl cyclohexylalanine inhibitor

Release Date: 06-Nov-2007 Exp. Method:

Characteristics Resolution: 2.10 Å

Hydrolase

Polymer: 1 Molecule: Dipeptidyl peptida

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	Compound	Mutation: T39S Chains: A,B EC no.: 2., Polymer: 2 Molecule: SUGAR (2-MER) Polymer: 4 Molecule: SUDIUM ION Polymer: 5 Molecule: SOJIUM ION PLUORIDYPRROLIDIN: 1-YLI-N,N-DIN OXO-2-(TRANS-4-I1,2,4)TRIAZOLOJI 5-YLOYCLOHEXYL) BUTANAMIDE
	Authors	Scapin, G.
☑ 2qib		Human dipeptidyl peptidase tv/ cd26 in complex with a 4-atyl cyclohexylalanine inhibitor
	Characteristics	Release Date: 06-Nov-2007 Exp. Method:
J. 18 10 10 10 10 10 10 10 10 10 10 10 10 10		Resolution: 2.25 Å
A. 16.	Classification	Hydrolase
	Compound	Polymer: 1 Moterule: Dipeptidyl peptida Mutation: T39S Orbiner A,B G R C no.: 3 , r Polymer: 2 Moterule: SUGAR (?-MER) Polymer: 4 Moterule: SODIUM (ON Polymer: 5 Moterule: (SS,SY-3-AMINO DIFLUOROPYRROLIDIN-1-Y)-N,N-L OXO-2 (TRANS-4-[1,2,4]TRIAZOLO[1 6-YLCYCLOHEXYL)BUTANAMIDE
	Authors	Scapin, G.
₹ 3c43		Human dipaptidyl paptidase IV/ CD28 in complex with a flouroplefin inhibitor
124	Characteristics	Release Date: 22-Apr-2008 Exp. Method:
AK. C.	Characteristics	Resolution: 2.30 Å
100	Classification	Hydrolase
	Compound	Polymer: 1 Molecule: Dipeptidyl peptida Fragmen: Catalytic domain Mutation: S Chains: A,B EC no.: 3.4.14.5 Polymer: 2 Molecule: SUGAR (2-MER) Polymer: 4 Molecule: SUGAR (2-MER) Polymer: 5 Molecule: SUGAR (2-MER)
	Authors	Scapin, G., Edmondson, S.D., Webs
₹ 3645		Human dipeptidyl peptidase tV/CD26 in complex with a fluorpoletin inhibitor
12 80	Characteristics	Release Date. 22-Apr-2008 Exp. Method.
	Characteristics	Resolution: 2.05 Å
	Classification	Hydrolase
	Compound	Polymer 1 Molecule: Dipeptidyl peptida Fragment: Catalytic domain Matation S Chains: A,B EC no. 3 4,14.5 © Polymer: 2 Molecule: SUGAR (2-MER) Polymer: 4 Molecule: SUGAR (2-MER)
	Authors	Scapin, G., Edmondson, S.D., Webs

	₽ 2178		Crystal structure of human dipeptidyl peptidses (V (DPP IV) complexed with ABT 341 a cyclohexene-constrained phenethylamine inhibitor
	Characteristics	Release Date: 09-Oct-2007 Exp. Method:	
		Resolution: 2.50 Å	
	Classification	Hydrolase	
		Compound	Polymer: 1 Molecule: Dipeptidyl peptida Chains: A,B,C,D EC no.: 3.4.14.5
		Authors	Longenecker, K.L., Pei, Z., Li, X.
	₱ 2hha		The structure of DPP4 in complex with an exadiazole inhibitor
		011	Release Date: 19-Sep-2006 Exp. Method.
	Characteristics	Resolution: 2.35 Å	
	Classification	Hydrolase	
			Polymer: 1 Molecule: Hypothetical prote Fragment: catalytic domain (39-766) M
		Compound	Chains: A,B EC no.: 3.4.14.5 Polymer 2 Molecule: SUGAR (2-MER) Polymer: 4 Molecule: SUGAR (2-MER) Polymer 5 Molecule: SUGAR (2-MER)
		Authors	Scapin, G.
	™ 2z3w		Prolyl tripeptidyl aminopeptidase mutani E636A
	A153.	Characteristics	Release Date: 19-Feb-2008 Exp. Method:
	Cirar acrorratios	Resolution: 2.00 Å	
	Classification	Hydrolase	
		Compound	Polymer: 1 Molecule: Dipeptidyl aminop Fragmeni: residues 39-732 Mutation: Et Chains A EC no.: 3.4.14
		Authors	Xu, Y., Nakajima, Y., ito. K., Yoshi
	™ 2z3z		Prolyl tripeptidyl aminopeptidase mutant E636A complexd with an inhibitor
	Characteristics	Release Date: 19-Feb-2008 Exp. Method:	
		Resolution: 1.95 Å	
	Classification	Hydrolase	
	The Marie		
		Compound	Polymer: 1 Motecule: Dipeptidyl aminop Fragment: residues 39-732 Mutation: Et Chains: A EC no.: 3.4.14.
		Compound Authors	Fragment: residues 39-732 Mutation: Et

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